

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
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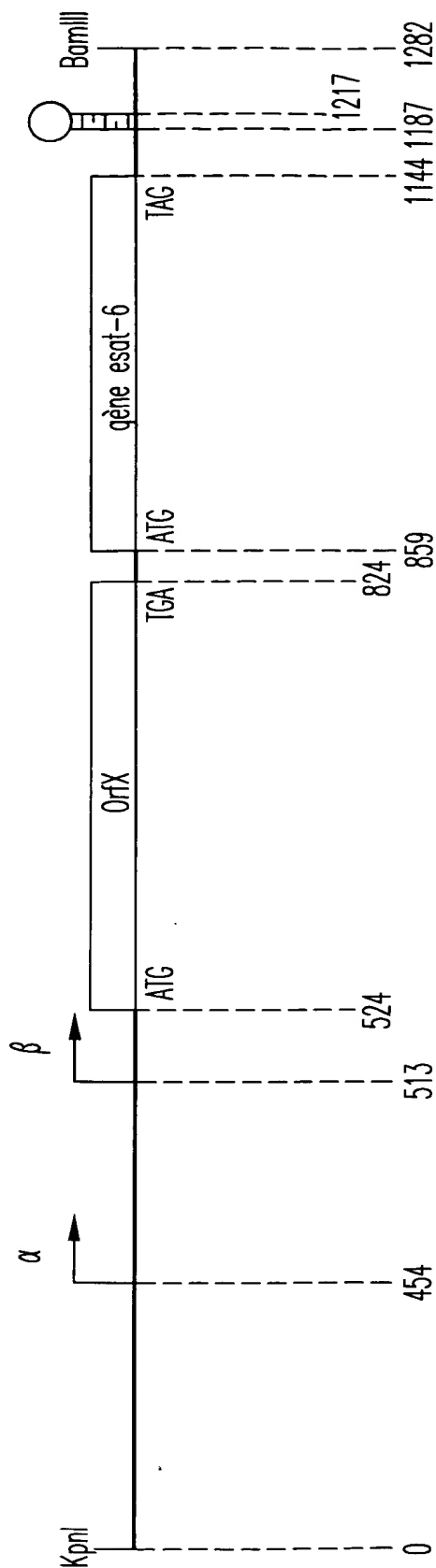


FIG. 1

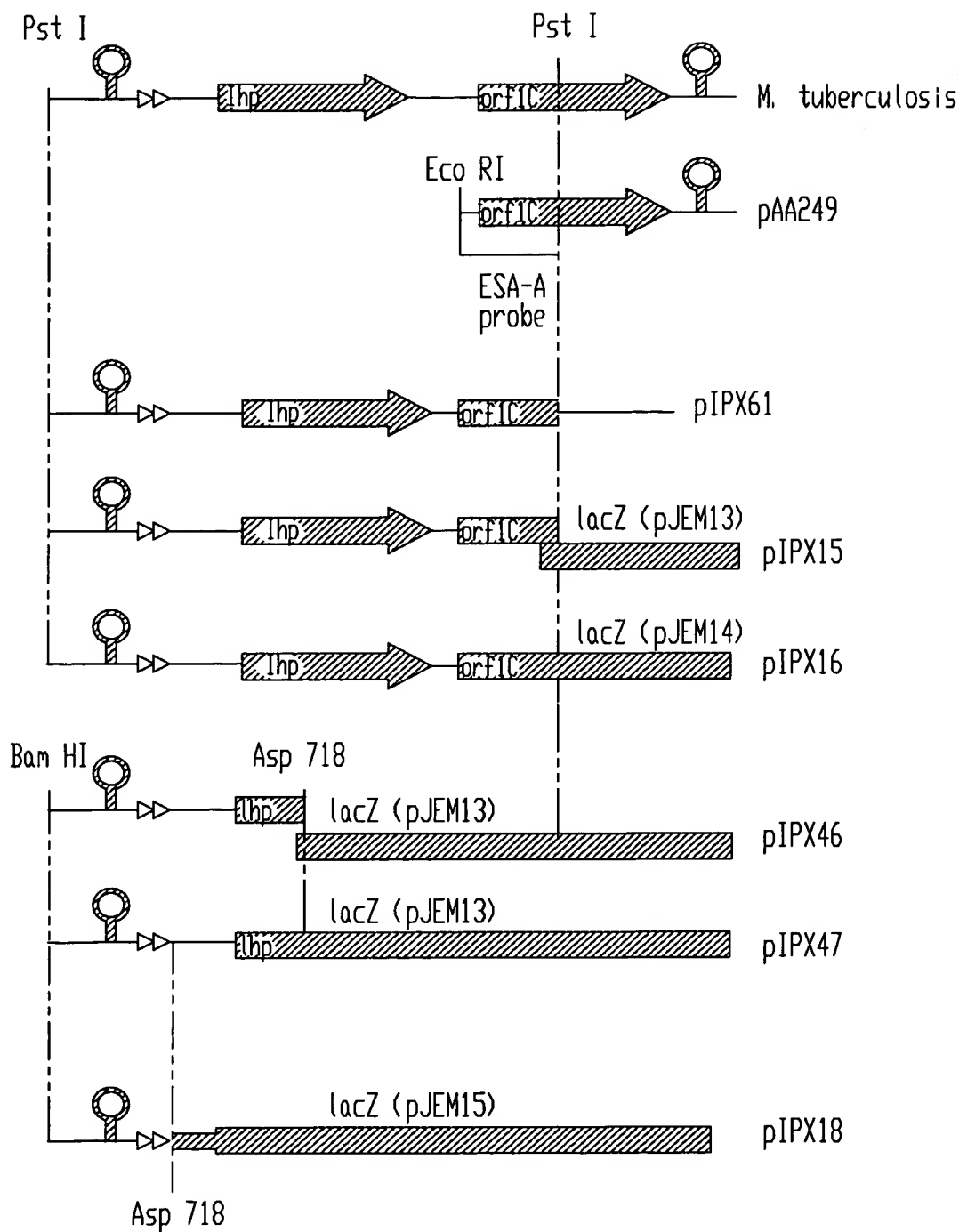


FIG. 2

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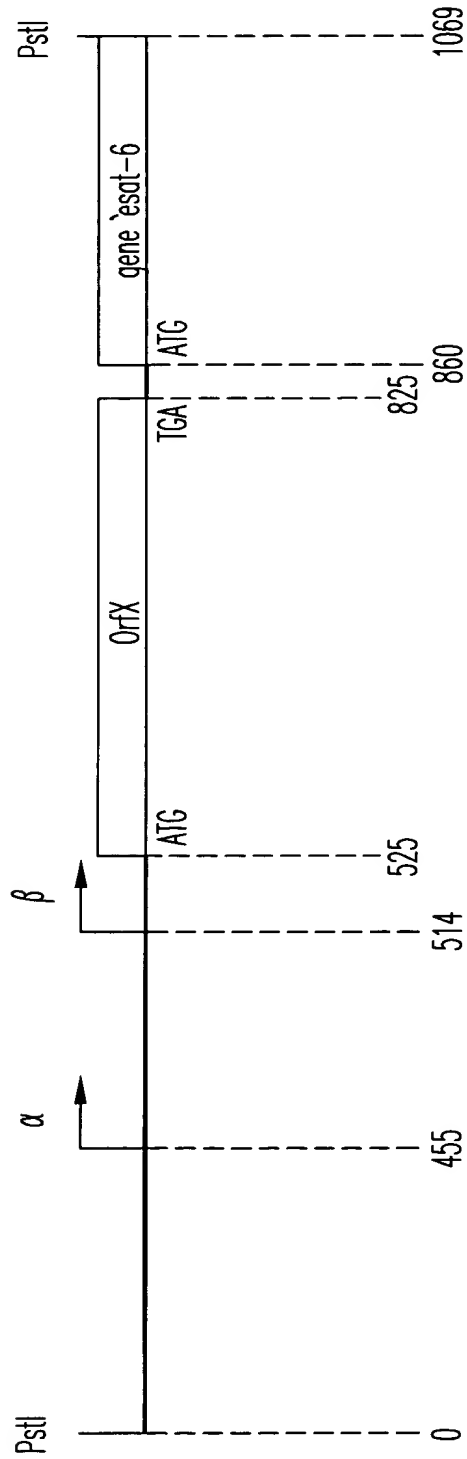
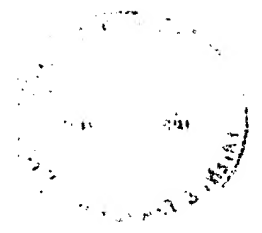


FIG. 3



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10 30 50 70
PstI
CTGCAGCAGGTGACGTCGTTGTTTCAGCCAGGTGGGCGGCACCGGCGGCGGCAACCCAGCCGACGAGGAAGCCGCGCAGAT
90 110 130 150
GGGCCTGCTCGGCACCAGTCCGCTGTCGAACCATCCGCTGGCTGGTGGATCAGGCCCCAGCGCGGGCGCGGGCCTGCTGC
170 190 210 230
GCGCGGAGTCGCTACCTGGCGCAGGTGGGTGCTTGACCCGCACGCCGCTGATGTCTCAGCTGATCGAAAAGCCGGTTGCC
250 270 290 310
CCCTCGGTGATGCCGGCGGCTGTTGCCGGATCGTCGGTGACGGGTGGCGCCGCTCCGGTGGGTCCGGGAGCGATGGGCCA
330 350 370 390
GGGTTGCAATCCGGCGGCTCCACCAGCCCGGGTCTGGTCGCGCCGGCACCGCTCGCGCAGGAGCGTGAAGAAGACGACG
410 430 450 470
AGGACGACTGGGACGAAGAGGACGACTGGTGAGCTCCCGTAATGACAACAGAC^{+1 Mtb.}TCCCGGCCACCCGGGCCGAAGACTT
490 510 530 550
GCCAACATTTTGGCGAGGAAGGTAAAGAGAGA^{+1 Ms}AGTAGTCCAGCATGGCAGAGATGAAGACCGATGCCGCTACCCTCGGG
570 590 610 630
CAGGAGGCAGGTAATTTTCGAGCGGATCTCCGGCGACCTGAAAACCCAGATCGACCAGGTGGAGTCGACGGCAGGTTTCGTT
Q E A G N F E R I S G D L K T Q I D Q V E S T A G S L
650 670 690 710
GCAGGGCCAGTGGCGCGGCGCGGCGGGGACGGCCGCCAGGCCGCGGTGGTGGCTTCCAAGAAGCAGCCAATAAGCAGA
Q G Q W R G A A G T A A Q A A V V R F Q E A A N K Q K
730 750 770 790
AGCAGGAACCTCGACGAGATCTCGACGAATATTCGTCAGGCCGGCGTCCAATACTCGAGGGCCGACGAGGAGCAGCAGCAG
Q E L D E I S T N I R Q A G V Q Y S R A D E E Q Q Q
810 830 850
esat-6
GCGCTGTCTCGCAAATGGGCTTCTGACCCGCTAATACGAAAAGAAACGGAGCAAAAACATGACAGAG.
A L S S Q M G F * M T E

FIG. 4A

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GATC12

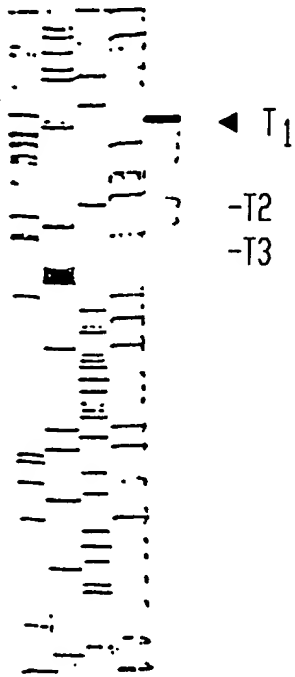


FIG. 5A

GATC12

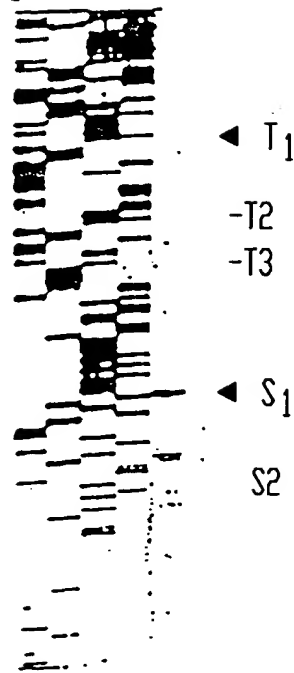


FIG. 5C

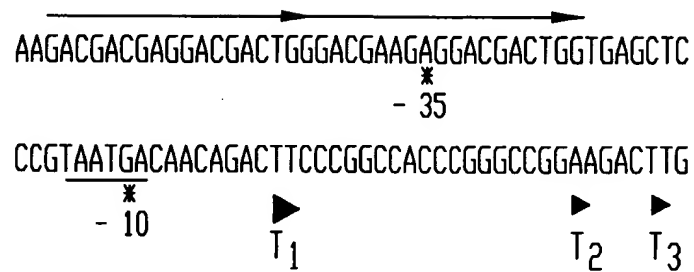


FIG. 5B

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1 2 3 4 5

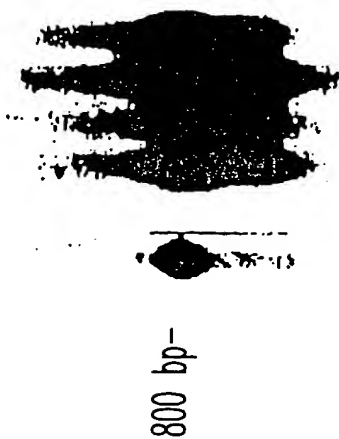
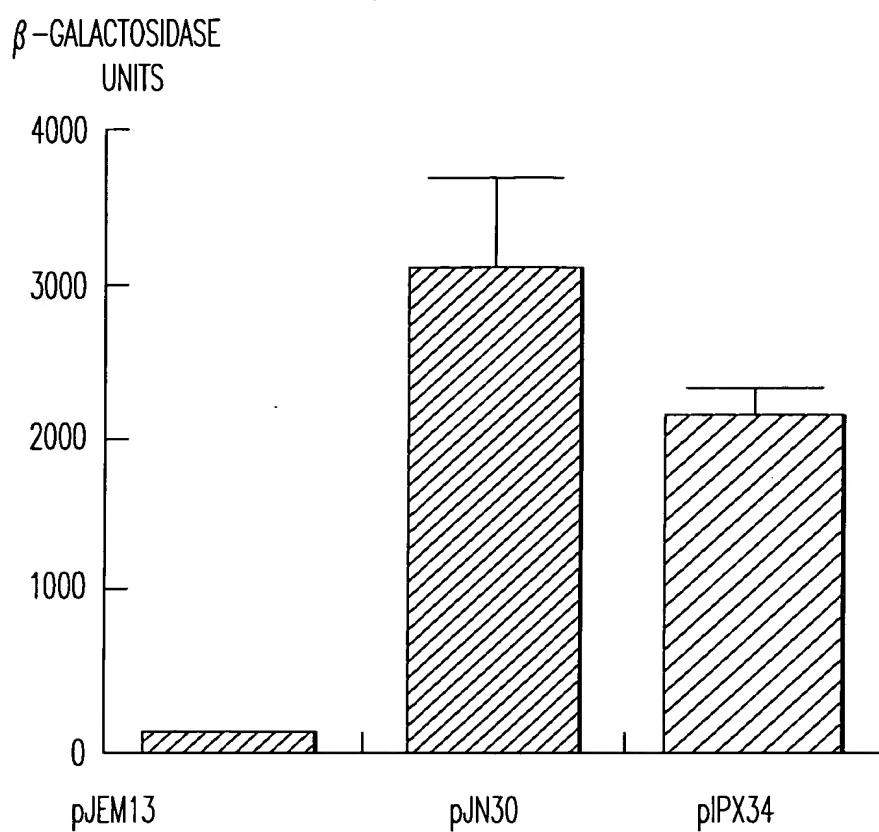


FIG. 6



ESAT-6 transcription terminator

FIG. 7

*FIG. 8*

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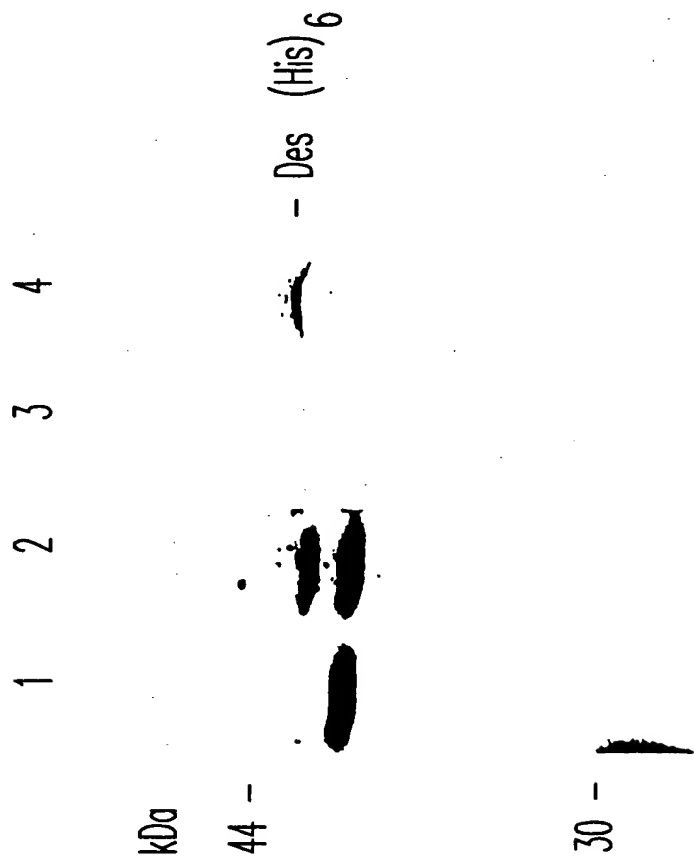


FIG. 9

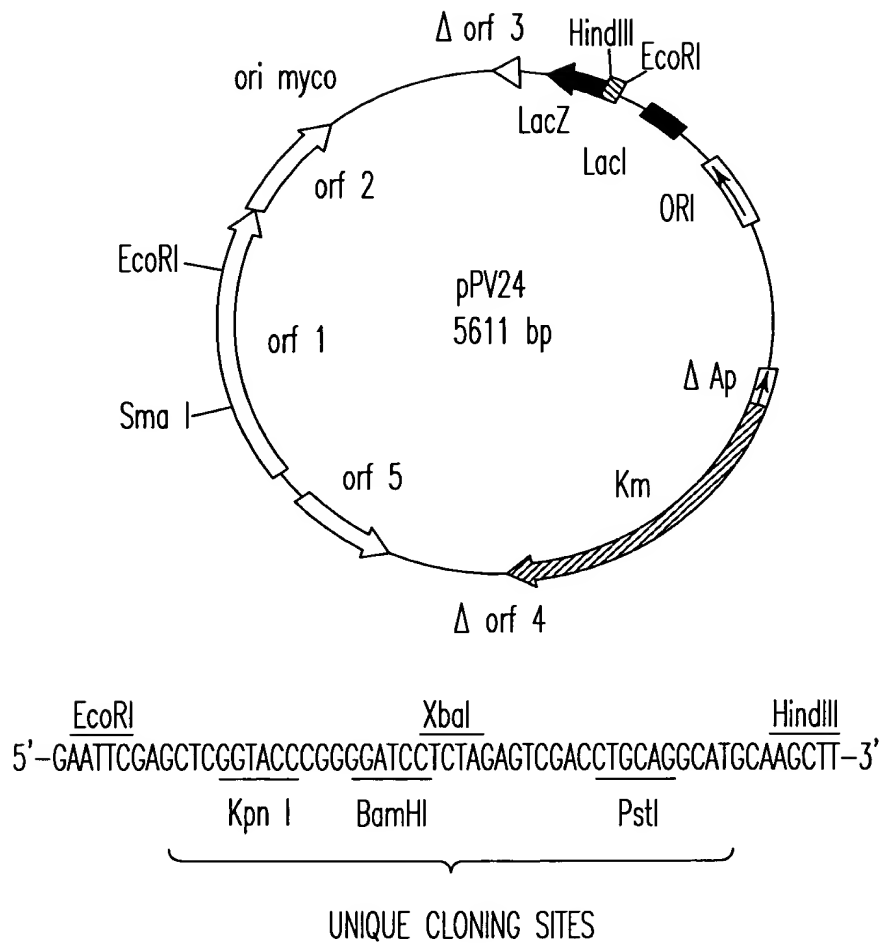


FIG. 10

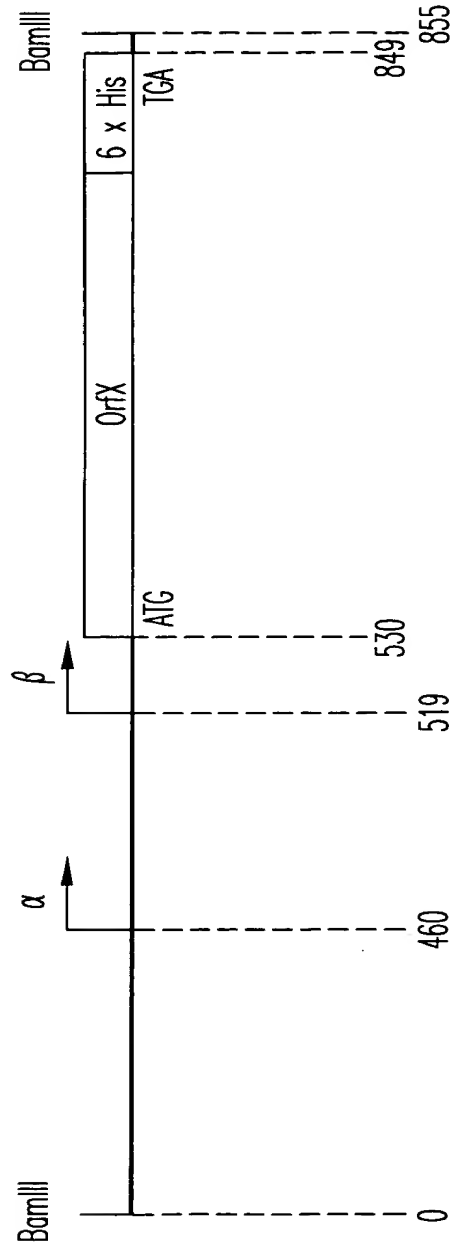


FIG. 11

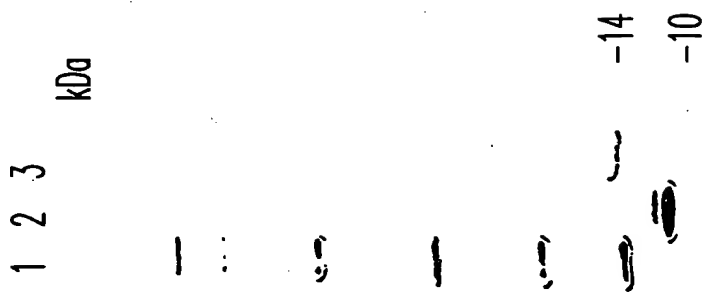


FIG. 12B



FIG. 12A

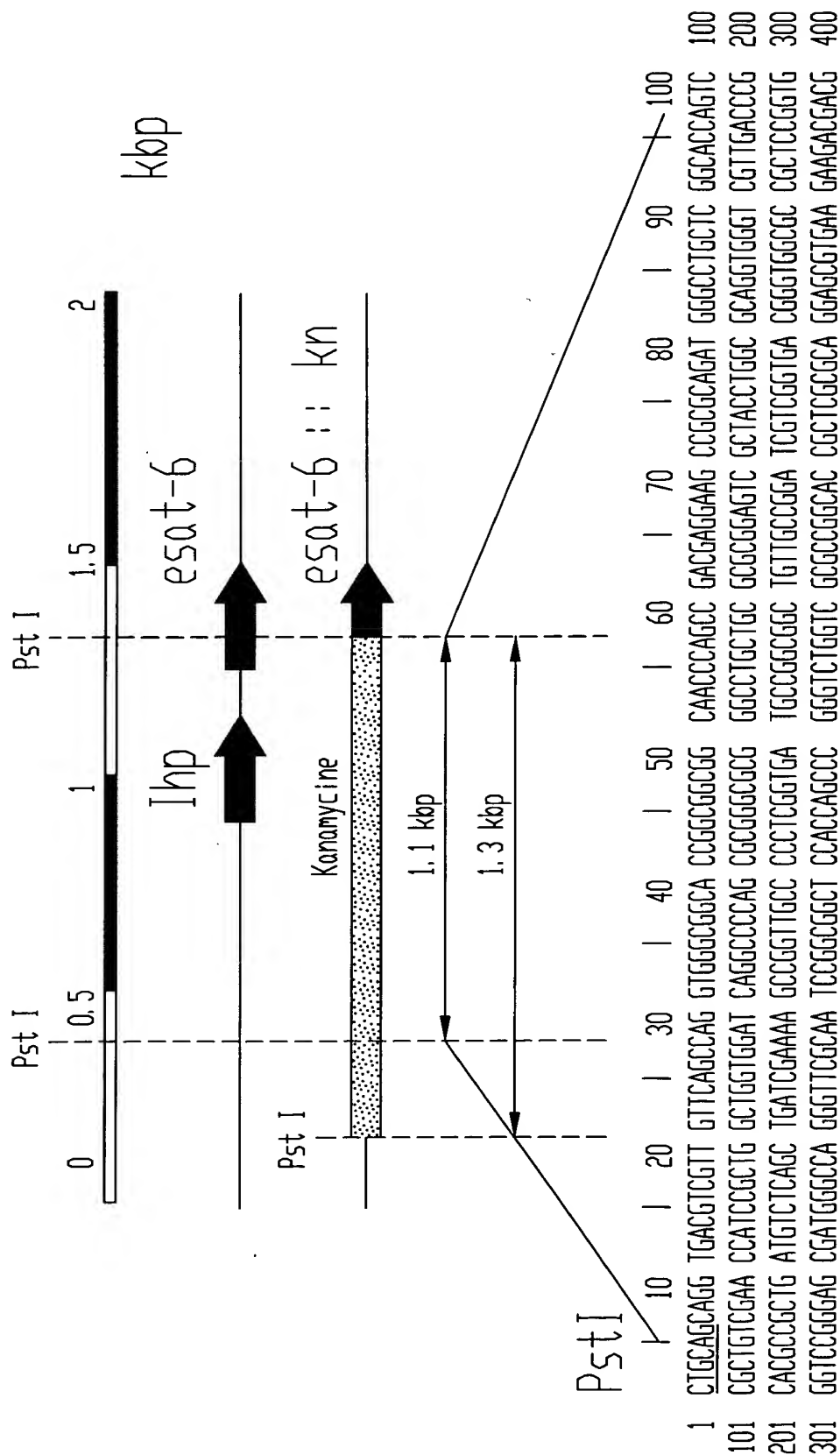


FIG. 13A

APPROVED	C. G. F.	OBLON ET AL (703)
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401	AGGACGACTG	GGACGAAGAG	GACGACTGGT	GAGCTCCCGT	AATGACAACA	GACTTCCCGG	CCACCCGGGC	CGGAGACTT	GCCAACATTT	TGGCGAGGAA	500
501	GGTAAGAGA	GAAAGTAGTC	CAGCATGGCA	GAGATGAGA	CCGATGCCGC	TACCCTCGGG	CAGGAGGCAG	GTAATTCGA	GGGATCTCC	GGCGACCTGA	600
601	AAACCCAGAT	CGACCAAGTG	GAGTCGACGG	CAGGTTTCGT	GCAGGGCCAG	TGGCGGGGG	CGGCGGGGAC	GGCGGCCAG	GCCGCGGTGG	TGCGCTTCCA	700
701	AGAGGACGCC	AATAAGCAGA	AGCAGGAAT	CGACGAGATC	TCGACGAATA	TTCGTGAGGC	CGGCGTCCAA	TACTCGAGGG	CCGACGAGGA	GCAGCAGCAG	800
801	GCGCTGTCT	CGCAATGGG	CTTCTGACCC	GCTAATACCA	AAAGAAACGG	AGCAAAACA	TCACAGAGCA	GCAGTGGAT	TTCGCGGGTA	TCGAGGCCGC	900
901	GGCAAGGCCA	ATCCAGGGAA	ATGTCAGTC	CATTCAATCC	CTCCTTGACG	AGGGGAAGCA	GTCCCTGACC	AAGCTCCAG	CGGCTGGGG	CGGTAGCGGT	1000
1001	TCGAGGCGT	ACCAGGGTGT	CCAGCAAAA	TGGGACGCCA	CGGCTACCGA	GCTGAACAAC	GGCCTGCAG				1069

FIG. 13B PstI